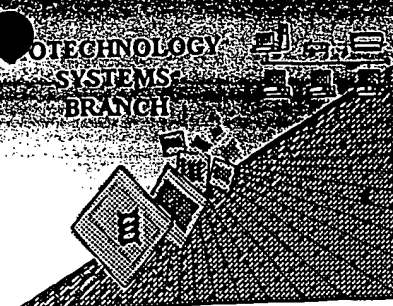
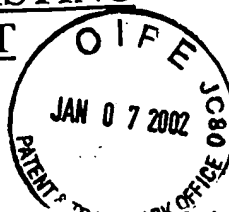


RAW SEQUENCE LISTING
ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/804014

Source: O I P E

Date Processed by STIC: 09/19/01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be downloaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>



OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/804,014

DATE: 09/19/2001
TIME: 13:53:55

Input Set : A:\Cura-221.app
Output Set: N:\CRF3\09192001\I804014.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: Li, Li
4 Padigaru, Muralidhara
5 Vernet, Corine
6 Fernandes, Elma
7 Shimkets, Richard
8 Spaderna, Steven
9 Majumder, Kumud
11 <120> TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoding Same
13 <130> FILE REFERENCE: 15966-721 US
15 <140> CURRENT APPLICATION NUMBER: 09/804,014
C--> 16 <141> CURRENT FILING DATE: 2001-09-07
18 <150> PRIOR APPLICATION NUMBER: 60/188,316
19 <151> PRIOR FILING DATE: 2000-03-10
21 <150> PRIOR APPLICATION NUMBER: 60/188,277
22 <151> PRIOR FILING DATE: 2000-03-10
24 <150> PRIOR APPLICATION NUMBER: 60/189,139
25 <151> PRIOR FILING DATE: 2000-03-14
27 <150> PRIOR APPLICATION NUMBER: 60/189,140
28 <151> PRIOR FILING DATE: 2000-03-14
30 <150> PRIOR APPLICATION NUMBER: 60/190,401
31 <151> PRIOR FILING DATE: 2000-03-17
33 <150> PRIOR APPLICATION NUMBER: 60/190,231
34 <151> PRIOR FILING DATE: 2000-03-17
36 <160> NUMBER OF SEQ ID NOS: 75
38 <170> SOFTWARE: PatentIn Ver. 2.1

Errored
Sequence indicates
1,949 nucleotides in
sequence "one". Found
only 594 nucleotides
in the sequence

ERRORED SEQUENCES

40 <210> SEQ ID NO: 1
41 <211> LENGTH: 1949
42 <212> TYPE: DNA
43 <213> ORGANISM: Homo sapiens
45 <400> SEQUENCE: 1
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47 ctgggaactc tggcaccttg ggctgtggaa ggctctggaa agtccttcaa agctggagtc 120
48 tgtcctccta agaaatctgc ccagtgcctt agatacaaga aacctgagtg ccagagtgc 180
49 tggcagtgtc caggggaagaa gagatgttgt cctgacactt gtggcatcaa atgcctggat 240
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53 ctgccatatg gaggaggctc tggagtcctg ctctgtgtgg tccaggctct tccaccctg 480
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/804,014

DATE: 09/19/2001

TIME: 13:53:56

Input Set : A:\Cura-221.app

Output Set: N:\CRF3\09192001\I804014.raw

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L:55 M:252 E: No. of Seq. differs, <211>LENGTH:Input:1949 Found:594 SEQ:1
L:2096 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40